

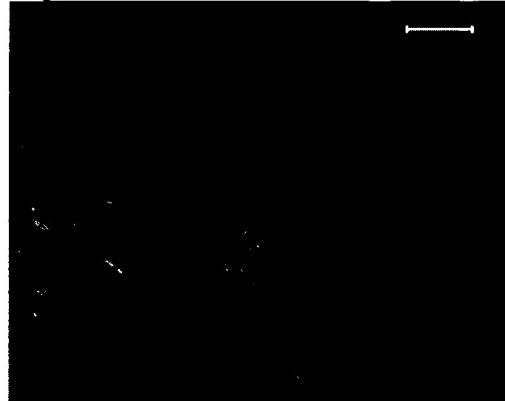
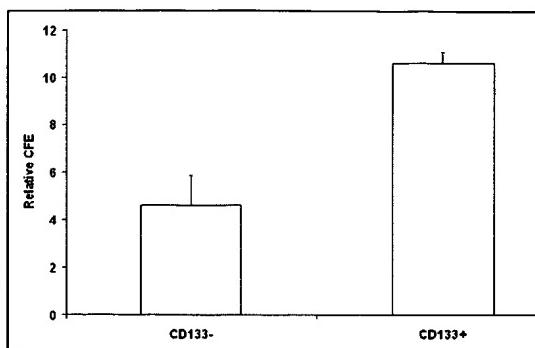
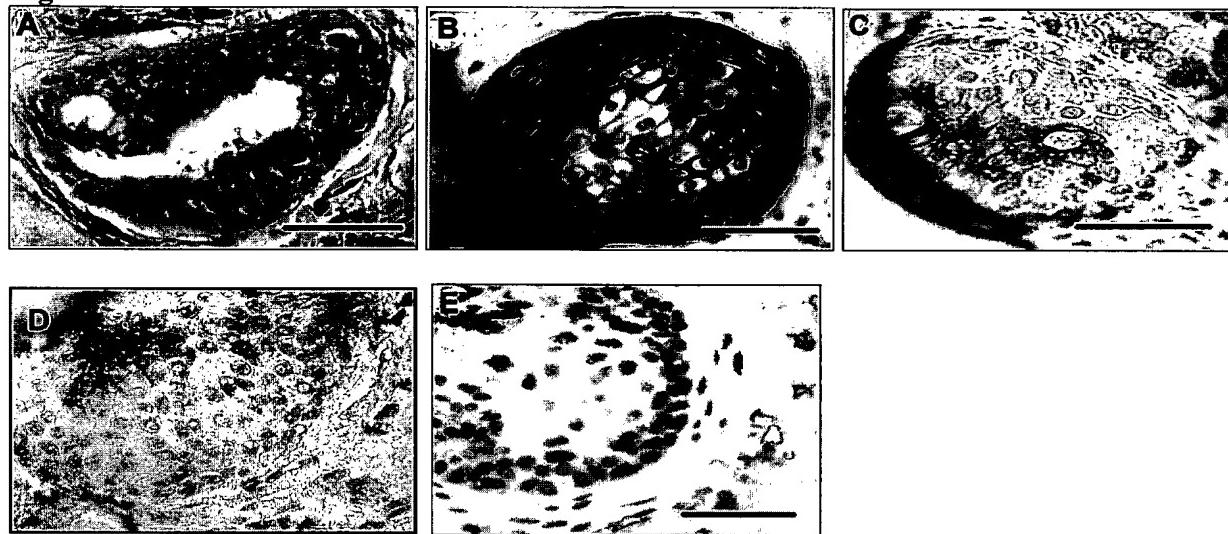
Fig 1A**Fig 1B****Fig 1C**

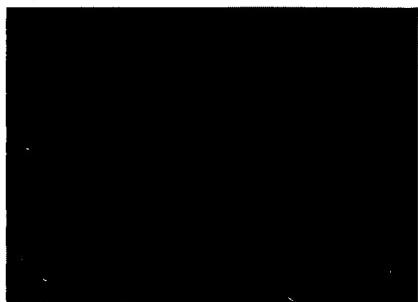
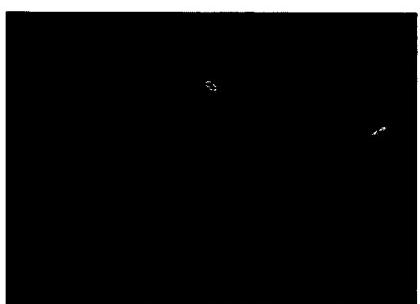
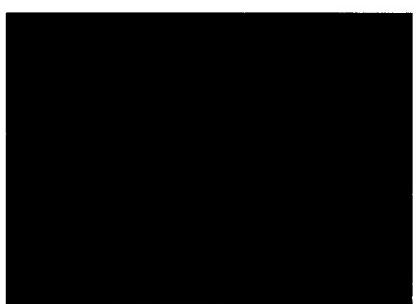
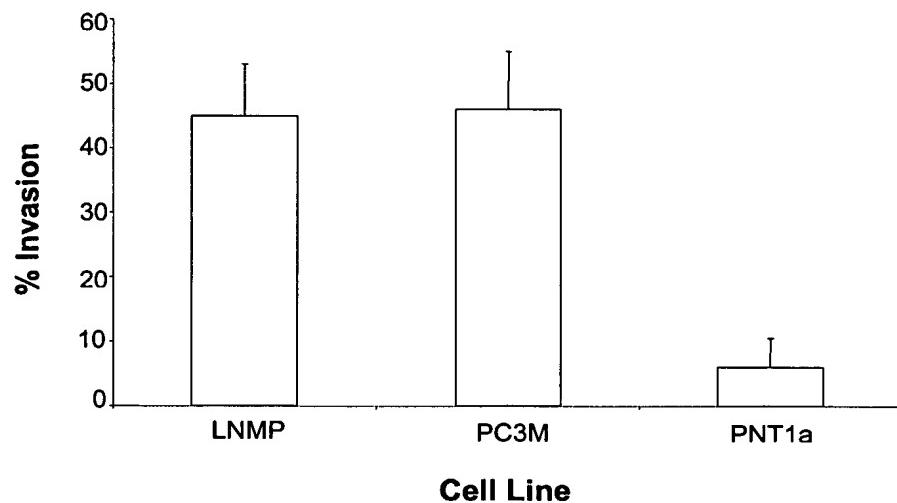
Fig 2A**2/4****Vimentin****Cytokeratin 18****E-Cadherin****PSA****Fig 2B**

Figure 3

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MHRTTRIKITELNPHLMCVLCGGYFIDATTIECLHSFKTCIV
RYLETSKYCPICDVQVHKTRPLLNIRSDKTLQDIVYKLVPGLFKNEMKRRDFYAAHP
SADAANGSNEDRGEVADEDKRIITDDEIISLSIEFFDQNRLDRKVNKDKEKSKEEVND
KRYLRCPAAMTVMHLRKFLRSKMDIPNTFQIDVMYEEPLKDYYTLMDIAYIYTWRN
GPLPLKYRVRPTCKRMKISHQRDGLTNAGELESDGSDKANSPAGGIPOSTSSCLPSPSTPVQ
SPHPQFPHISSTMNGTSNSPSGNHQSSFANPRKSSVNGSSATSSG

Figure 4

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1 CAGCAACTAT GAAATAATCG TAGTATGAGA GGCAGAGATC GGGCGAGAC AATGGGGATG
 61 TGGGCGCGGG AGCCCCGTTG CGGCTTAGCA GCACCTCCCA GCCCGCGAGA ATAAAACCGA
 121 TCGCCGCCCG TCCGCGCGCG CCCTCCCCG AGTGCAGGAGC GGGAGGAGGC GGCGCGGGCC
 181 GAGGAGGAGG AGGAGGAGGC CCCGGAGGAG GAGGCGTTGG AGGTCGAGGC GGAGGCGGAG
 241 GAGGAGGAGG CCGAGGCGCC GGAGGAGGCC GAGGCGCCGG AGCAGGAGGA GGCGGGCCGG
 301 AGGCGGCATG AGACGAGCGT GGCGGCCGCG GCTGCTCGGG GCCGCGCTGG TTGCCCATTG
 361 ACAGCGCGT CTGCAGCTCG CTTCAAGATG GCCGCTTGGC TCGCATTATCAT TTTCTGCTGA
 421 ACGACTTTA ACTTTCATTG TCTTTCCGC CGCCTTCGAT CGCCTCGC CGGCTGCTCT
 481 TTCCGGGATT TTTTATCAAG CAGAACATGCA TCGAACAAACG AGAACATCAAAGA TCACTGAGCT
 541 AAATCCCCAC CTGATGTGT TGCTTGTGG AGGGTACTTC ATTGATGCCA CAACCATAAT
 601 AGAATGCTA CATTCCCTTC GTAAAACGTTG TATTGTTCTG TACCTGGAGA CCAGCAAGTA
 661 TTGTCCTATT TGTGATGTCC AAGTTCACAA GACCAGACCA CTACTGAATA TAAGGTCAAGA
 721 TAAAACCTCTC CAAGATATTG TATACAAATT AGTCCAGGG CTTTTCAAAA ATGAAATGAA
 781 GAGAAGAAGG GATTTTATG CAGCTCATCC TTCTGCTGAT GCTGCCAATG GCTCTAATGA
 841 AGATAGAGGA GAGGTTGCAG ATGAAGATAA GAGAATTATA ACTGATGATG AGATAATAAG
 901 CTTATCCATT GAATTCTTG ACCAGAACAG ATTGGATCGG AAAGTAAACA AAGACAAAGA
 961 GAAATCTAAG GAGGAGGTGA ATGATAAAAG ATACTTACGA TGCCCAGCAG CAATGACTGT
 1021 GATGCACTTA AGAAAGTTTC TCAGAAGTAA AATGGACATA CCTAATACTT TCCAGATTGA
 1081 TGTCATGTAT GAGGAGGAAC CTTTAAAGGA TTATTATACA CTAATGGATA TTGCCTACAT
 1141 TTATACCTGG AGAAGGAATG GTCCACTTCC ATTGAAATAC AGAGTTCGAC CTACTTGTAA
 1201 AAGAATGAAG ATCAGTCACC AGAGAGATGG ACTGACAAAT GCTGGAGAAC TGAAAAGTGA
 1261 CTCTGGGAGT GACAAGGCCA ACAGCCCAGC AGGAGGTATT CCCTCCACCT CTTCTGTTT
 1321 GCCTAGCCCC AGTACTCCAG TGCACTCTCC TCATCCACAG TTTCCTCACAA TTTCCAGTAC
 1381 TATGAATGGA ACCAGCAACA GCCCCAGCGG TAACCACCAA TCTTCTTTG CCAATAGACC
 1441 TCGAAAATCA TCAGTAAATG GGTCACTCAGC AACTTCTTCT GGTGATACC TGAGACTGTT
 1501 AAGGAAAAAA ATTTAAACCC CCTGATTAT ATAGATATCT TCATGCCATT ACAGCTTTCT
 1561 AGATGCTAAT ACATGTGACT ATCGTCAAT TTGCTTCTT TTGTTAGTGAC ATAAATTTG
 1621 GCTATAAAAG ATGGACTACA TGTGATACTC CTATGGACGT TAATTGAAAA GAAAGATTG
 1681 TGTATATAAAG ATTGGTTTC TTGGAAAGCA GGCAGACTT TTTCTCTGTG TTAGGAAAGA
 1741 TGGGAAATGG TTTCTGTAAC CATTGTTGG ATTGGAAGT ACTCTGCACT GGACATAAGC
 1801 ATTGGCCAT AGTTGTTAA TCTCAACTAA CGCCTACATT ACATTCTCT TGATCGTTCT
 1861 TGTTATTACG CTGTTTGTG AACCTGTAGA AAACAGTGC TTCTTATCTT GAAATTCAAC
 1921 CAACGGAAAG AATATGCATA GAATAATGCA TTCTATGTC CCATGTCACT GTGAATAACG
 1981 ATTTCTGCA TATTTAGCCA TTTTGATTCC TTGTTGATT ATACTTCTCT GTTGCTACGC
 2041 AAAACCGATC AAAGAAAAAGT GAACTTCAGT TTACAATCT GTATGCTAA AAGCGGGTAC
 2101 TACCGTTTAT TTTACTGACT TGTTAAATG ATTCGTTTT GTAAGAATCA GATGGCATTAA
 2161 TGCTTGTGTTG ACAATGCCAT ATTGGTATAT GACATAACAG GAAACAGTAT TGTATGATAT
 2221 ATTTATAAAT GCTATAAAAGA AATATTGTGT TTCAATGCAATT CAGAAATGAT TGTTAAAATT
 2281 CTCCCAACTG GTTCGACCTT TGCAAGATACC CATAACCTAT GTTGAGCCTT GCTTACCAAGC
 2341 AAAGAATATT TTTAATGTGG ATATCTAATT CTAAGTCTG TTCCATTAGA AGCAATTGGC
 2401 ACATCTTCT ATACTTTTACTA TACTTTCTC CAGTAATACA TGTTTACTTT AAAAATTGTT
 2461 GCAGTGAAGA AAAACCTTTA ACTGAGAAAT ATGGAAACCG TCTTAATTCC CATTGGCTA
 2521 TGATGGAATT AATATTGTAT TTAAATGCAATT CATATTGTC ACTATAATTTC TAAAACAATT
 2581 TTAAATAAACCAGCAGTG TGCTAAAAGA AGGCATTAA TCTAAAGTTA TTAAATAGG
 2641 TGGTATAGCA GTAATTAAAT TTAAAGAGT TGCTTTTACA GTTAACAATG GAATATGCCT
 2701 TCTCTGCTAT GTCTGAAAT AGAAGCTATT TATTATGAGC TTCTACAGGT ATTTTTAAAT
 2761 AGAGCAAGCA TGTTGAATT AAAATATGAA TAACCCACC CAACAATTTC CAGTTTATTT
 2821 TTTGTTTGG TCGAACCTGG TGTGTTCA TCACCCATCA GTTATTGAGG AGGGTGTAA
 2881 TTCTATATGA ATATTGTTTCA ATGTTGTTAT GGGAAAATTG TAGCTAAACA TTTCATTGTC
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 3181 AAAAGGATA TTCATTTAGA AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA
 3241 AAAAAAAAAA A